# <u>RAW SEQUENCE LISTING</u>

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Application Serial Number: _____ 10/578, 139 _____

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**RAW SEQUENCE LISTING**          DATE: 03/01/2007
**PATENT APPLICATION:   US/10/578,139**          TIME: 14:45:41

Input Set : **N:\efs\03_01_07\10578139_efs\2007022725200132PUS1ST25.txt**
Output Set: **N:\CRF4\03012007\J578139.raw**

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 3 <110> APPLICANT: MIYAGAWA , Shuji
 4        MATSUNAMI , Katsuyoshi
 6 <120> TITLE OF INVENTION: HLA-E CHIMERIC MOLECULE
 8 <130> FILE REFERENCE: 2520-0132PUS1
10 <140> CURRENT APPLICATION NUMBER: US 10/578,139
11 <141> CURRENT FILING DATE: 2006-05-03
13 <160> NUMBER OF SEQ ID NOS: 92
15 <170> SOFTWARE: PatentIn version 3.4
17 <210> SEQ ID NO: 1
18 <211> LENGTH: 21
19 <212> TYPE: PRT
20 <213> ORGANISM: Artificial Sequence
22 <220> FEATURE:
23 <223> OTHER INFORMATION: Description of Artificial Sequence: Synthetic chimeric
sequence
24        SP of HLA-E
27 <400> SEQUENCE: 1
29 Met Val Asp Gly Thr Leu Leu Leu Leu Ser Glu Ala Leu Ala Leu
30 1              5                  10                 15
33 Thr Gln Thr Trp Ala
34              20
37 <210> SEQ ID NO: 2
38 <211> LENGTH: 90
39 <212> TYPE: PRT
40 <213> ORGANISM: Artificial Sequence
42 <220> FEATURE:
43 <223> OTHER INFORMATION: Description of Artificial Sequence: Synthetic chimeric
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44        a1 domain of HLA-E
47 <400> SEQUENCE: 2
49 Gly Ser His·Ser Leu Lys Tyr Phe His Thr Ser Val Ser Arg Pro Gly
50 1              5                  10                 15
53 Arg Gly Glu Pro Arg Phe Ile Ser Val Gly Tyr Val Asp Asp Thr Gln
54              20                 25                 30
57 Phe Val Arg Phe Asp Asn Asp Ala Ala Ser Pro Arg Met Val Pro Arg
58         35                 40                 45
61 Ala Pro Trp Met Glu Gln Glu Gly Ser Glu Tyr Trp Asp Arg Glu Thr
62     50                 55                 60
65 Arg Ser Ala Arg Asp Thr Ala Gln Ile Phe Arg Val Asn Leu Arg Thr
66 65                 70                 75                 80
69 Leu Arg Gly Tyr Tyr Asn Gln Ser Glu Ala
70              85                 90
73 <210> SEQ ID NO: 3
74 <211> LENGTH: 92
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75 <212> TYPE: PRT

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Output Set: N:\CRF4\03012007\J578139.raw

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 76 <213> ORGANISM: Artificial Sequence
 78 <220> FEATURE:
 79 <223> OTHER INFORMATION: Description of Artificial Sequence: Synthetic chimeric
sequence
 80       a2 domain of HLA-E
 83 <400> SEQUENCE: 3
 85 Gly Ser His Thr Leu Gln Trp Met His Gly Cys Glu Leu Gly Pro Asp
 86 1               5                  10                · 15
 89 Arg Arg Phe Leu Arg Gly Tyr Glu Gln Phe Ala Tyr Asp Gly Lys Asp
 90             20                  25                  30
 93 Tyr Leu Thr Leu Asn Glu Asp Leu Arg Ser Trp Thr Ala Val Asp Thr
 94         35                  40                  45
 97 Ala Ala Gln Ile Ser Glu Gln Lys Ser Asn Asp Ala Ser Glu Ala Glu
 98     50                  55                  60
101 His Gln Arg Ala Tyr Leu Glu Asp Thr Cys Val Glu Trp Leu His Lys
102 65                  70                  75              .  80
105 Tyr Leu Glu Lys Gly Lys Glu Thr Leu Leu His Leu
106                 85                  90
109 <210> SEQ ID NO: 4
110 <211> LENGTH: 92
111 <212> TYPE: PRT
112 <213> ORGANISM: Artificial Sequence
114 <220> FEATURE:
115 <223> OTHER INFORMATION: Description of Artificial Sequence: Synthetic chimeric
sequence
116       a3 domain of HLA-E
119 <400> SEQUENCE: 4
121 Glu Pro Pro Lys Thr His Val Thr His His Pro Ile Ser Asp His Glu
122 1               5                  10                  15
125 Ala Thr Leu Arg Cys Trp Ala Leu Gly Phe Tyr Pro Ala Glu Ile Thr
126             20                  25                  30
129 Leu Thr Trp Gln Gln Asp Gly Glu Gly His Thr Gln Asp Thr Glu Leu
130         35                  40                  45
133 Val Glu Thr Arg Pro Ala Gly Asp Gly Thr Phe Gln Lys Trp Ala Ala
134     50                  55                  60
137 Val Val Val Pro Ser Gly Glu Glu Gln Arg Tyr Thr Cys His Val Gln
138 65              70                  75                  80
141 His Glu Gly Leu Pro Glu Pro Val Thr Leu Arg Trp
142                 85                  90
145 <210> SEQ ID NO: 5
146 <211> LENGTH: 63
147 <212> TYPE: PRT
148 <213> ORGANISM: Artificial Sequence
150 <220> FEATURE:
151 <223> OTHER INFORMATION: Description of Artificial Sequence: Synthetic chimeric
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152       Transmembrane domain of HLA-E ·
155 <400> SEQUENCE: 5
157 Lys Pro Ala Ser Gln Pro Thr Ile Pro Ile Val Gly Ile Ile Ala Gly
158 1               5                  10                  15
161 Leu Val Leu Leu Gly Ser Val Val Ser Gly Ala Val Val Ala Ala Val
162             20                  25                  30
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165 Ile Trp Arg Lys Lys Ser Ser Gly Gly Lys Gly Gly Ser Tyr Ser Lys
166         35                  40                  45
169 Ala Glu Trp Ser Asp Ser Ala Gln Gly Ser Glu Ser His Ser Leu
170      50                  55                  60
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174 <211> LENGTH: 63
175 <212> TYPE: DNA
176 <213> ORGANISM: Artificial Sequence
178 <220> FEATURE:
179 <223> OTHER INFORMATION: Description of Artificial Sequence: Synthetic chimeric
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186 gcg                                                                   63
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190 <211> LENGTH: 270
191 <212> TYPE: DNA
192 <213> ORGANISM: Artificial Sequence
194 <220> FEATURE:
195 <223> OTHER INFORMATION: Description of Artificial Sequence: Synthetic chimeric
sequence
196        a1 domain of HLA-E
199 <400> SEQUENCE: 7
200 ggctcccact ccttgaagta tttccacact tccgtgtccc ggcccggccg cggggagccc     60
202 cgcttcatct ctgtgggcta cgtggacgac acccagttcg tgcgcttcga caacgacgcc    120
204 gcgagtccga ggatggtgcc gcgggcgccg tggatggagc aggaggggtc agagtattgg    180
206 gaccgggaga cacggagcgc cagggacacc gcacagattt tccgagtgaa tctgcggacg    240
208 ctgcgcggct actacaatca gagcgaggcc                                     270
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212 <211> LENGTH: 276
213 <212> TYPE: DNA
214 <213> ORGANISM: Artificial Sequence
216 <220> FEATURE:
217 <223> OTHER INFORMATION: Description of Artificial Sequence: Synthetic chimeric
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218        a2 domain of HLA-E
221 <400> SEQUENCE: 8
222 gggtctcaca ccctgcagtg gatgcatggc tgcgagctgg ggcccgacag gcgcttcctc     60
224 cgcgggtatg aacagttcgc ctacgacggc aaggattatc tcaccctgaa tgaggacctg    120
226 cgctcctgga ccgcggtgga cacggcggct cagatctccg agcaaaagtc aaatgatgcc    180
228 tctgaggcgg agcaccagag agcctacctg gaagacacat gcgtggagtg gctccacaaa    240
230 tacctggaga aggggaagga gacgctgctt cacctg                              276
233 <210> SEQ ID NO: 9
234 <211> LENGTH: 276
235 <212> TYPE: DNA
236 <213> ORGANISM: Artificial Sequence
238 <220> FEATURE:
239 <223> OTHER INFORMATION: Description of Artificial Sequence: Synthetic chimeric
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240        a3 domain of HLA-E
243 <400> SEQUENCE: 9
244 gagcccccaa agacacacgt gactcaccac cccatctctg accatgaggc caccctgagg     60
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RAW SEQUENCE LISTING     DATE: 03/01/2007
PATENT APPLICATION: US/10/578,139  TIME: 14:45:41

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    Output Set: **N:\CRF4\03012007\J578139.raw**

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246 tgctgggccc tgggcttcta ccctgcggag atcacactga cctggcagca ggatggggag   120
248 ggccataccc aggacacgga gctcgtggag accaggcctg caggggatgg aaccttccag   180
250 aagtgggcag ctgtggtggt gccttctgga gaggagcaga gatacacgtg ccatgtgcag   240
252 catgaggggc tacccgagcc cgtcaccctg agatgg                            276
255 <210> SEQ ID NO: 10
256 <211> LENGTH: 192
257 <212> TYPE: DNA
258 <213> ORGANISM: Artificial Sequence
260 <220> FEATURE:
261 <223> OTHER INFORMATION: Description of Artificial Sequence: Synthetic chimeric
sequence
262        Transmembrane domain of HLA-E
265 <400> SEQUENCE: 10
266 aagccggctt cccagcccac catccccatc gtgggcatca ttgctggcct ggttctcctt    60
268 ggatctgtgg tctctggagc tgtggttgct gctgtgatat ggaggaagaa gagctcaggt   120
270 ggaaaaggag ggagctactc taaggctgag tggagcgaca gtgcccaggg gtctgagtct   180
272 cacagcttgt aa                                                     192
275 <210> SEQ ID NO: 11
276 <211> LENGTH: 24
277 <212> TYPE: PRT
278 <213> ORGANISM: Artificial Sequence
280 <220> FEATURE:
281 <223> OTHER INFORMATION: Description of Artificial Sequence: Synthetic chimeric
sequence
282        SP of HLA-G1
285 <400> SEQUENCE: 11
287 Met Val Val Met Ala Pro Arg Thr Leu Phe Leu Leu Leu Ser Gly Ala
288 1               5                   10                  15
291 Leu Thr Leu Thr Glu Thr Trp Ala
292             20
295 <210> SEQ ID NO: 12
296 <211> LENGTH: 90
297 <212> TYPE: PRT
298 <213> ORGANISM: Artificial Sequence
300 <220> FEATURE:
301 <223> OTHER INFORMATION: Description of Artificial Sequence: Synthetic chimeric
sequence
302        a1 domain of HLA-G1
305 <400> SEQUENCE: 12
307 Gly Ser His Ser Met Arg Tyr Phe Ser Ala Ala Val Ser Arg Pro Gly
308 1               5                   10                  15
311 Arg Gly Glu Pro Arg Phe Ile Ala Met Gly Tyr Val Asp Asp Thr Gln
312             20                  25                  30
315 Phe Val Arg Phe Asp Ser Asp Ser Ala Cys Pro Arg Met Glu Pro Arg
316         35                  40                  45
319 Ala Pro Trp Val Glu Gln Glu Gly Pro Glu Tyr Trp Glu Glu Glu Thr
320     50                  55                  60
323 Arg Asn Thr Lys Ala His Ala Gln Thr Asp Arg Met Asn Leu Gln Thr
324 65                  70                  75                  80
327 Leu Arg Gly Tyr Tyr Asn Gln Ser Glu Ala
328             85                  90
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333 <212> TYPE: PRT
334 <213> ORGANISM: Artificial Sequence
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337 <223> OTHER INFORMATION: Description of Artificial Sequence: Synthetic chimeric
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338       a2 domain of HLA-G1
341 <400> SEQUENCE: 13
343 Ser Ser His Thr Leu Gln Trp Met Ile Gly Cys Asp Leu Gly Ser Asp
344 1               5                   10                  15
347 Gly Arg Leu Leu Arg Gly Tyr Glu Gln Tyr Ala Tyr Asp Gly Lys Asp
348             20                  25                  30
351 Tyr Leu Ala Leu Asn Glu Asp Leu Arg Ser Trp Thr Ala Ala Asp Thr
352         35                  40                  45
355 Ala Ala Gln Ile Ser Lys Arg Lys Cys Glu Ala Ala Asn Val Ala Glu
356     50              55                  60
359 Gln Arg Arg Ala Tyr Leu Glu Gly Thr Cys Val Glu Trp Leu His Arg
360 65                  70                  75                  80
363 Tyr Leu Glu Asn Gly Lys Glu Met Leu Gln Arg Ala
364                 85                  90
367 <210> SEQ ID NO: 14
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369 <212> TYPE: PRT
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372 <220> FEATURE:
373 <223> OTHER INFORMATION: Description of Artificial Sequence: Synthetic chimeric
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374       a3 domain of HLA-G1
377 <400> SEQUENCE: 14
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380 1               5                   10                  15
383 Ala Thr Leu Arg Cys Trp Ala Leu Gly Phe Tyr Pro Ala Glu Ile Ile
384             20                  25                  30
387 Leu Thr Trp Gln Arg Asp Gly Glu Asp Gln Thr Gln Asp Val Glu Leu
388         35                  40                  45
391 Val Glu Thr Arg Pro Ala Gly Asp Gly Thr Phe Gln Lys Trp Ala Ala
392     50              55                  60
395 Val Val Val Pro Ser Gly Glu Glu Gln Arg Tyr Thr Cys His Val Gln
396 65                  70                  75                  80
399 His Glu Gly Leu Pro Glu Pro Leu Met Leu Arg Trp
400                 85                  90
403 <210> SEQ ID NO: 15
404 <211> LENGTH: 40
405 <212> TYPE: PRT
406 <213> ORGANISM: Artificial Sequence
408 <220> FEATURE:
409 <223> OTHER INFORMATION: Description of Artificial Sequence: Synthetic chimeric
sequence
410       Transmembrane domain of HLA-G1
413 <400> SEQUENCE: 15
415 Lys Gln Ser Ser Leu Pro Thr Ile Pro Ile Met Gly Ile Val Ala Gly
416 1               5                   10                  15
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**VERIFICATION SUMMARY**                  DATE: 03/01/2007
PATENT APPLICATION:  **US/10/578,139**    TIME: 14:45:42

Input Set : **N:\efs\03_01_07\10578139_efs\2007022725200132PUS1ST25.txt**
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